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<110> COSTA e SILVA, OSWALDO DA
      BOHNERT, HANS J.
      THIELEN, NOCHA VAN
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Ser Ser Val Thr Glu Ile Lys Ile Gly Ser Arg Gly Leu Asn Gly Asn 85 90 95

Phe Asn Pro Ser Tyr Phe Gln Asn Ala Phe Lys Lys Leu Arg Ile Phe 100 105 110

Asp Ala Ser Asn Asn Ile Glu Gly Asn Ile Pro Gln Gln Phe Pro 115 120 125

Thr Ser Leu Thr Gln Met Ile Leu Asn Asn Lys Leu Thr Gly Gly 130 135 140

Leu Pro Gln Phe Asp Gln Leu Gly Ala Leu Thr Val Val Asn Leu Ser 145 150 155 160

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- Gln Asn Asn Lys Phe Asn Gly Lys Leu Pro Asp Asp Phe Ser Arg Leu 210 215 220
- Lys Asn Leu Gln Thr Phe Asn Ile Glu Asn Asp Gln Phe Thr Gly Asn 225 230 235 240
- Tyr Pro Ser Gly Leu Pro Ser Asn Ser Arg Val Gly Gly Asn Arg Leu 245 250 255
- Thr Phe Pro Pro Pro Pro Ala Pro Gly Thr Pro Ala Pro Arg Thr Pro 260 265 270
- Ser Pro Ser Gly Thr Ser Asn Gly Ser Ser Ser His Leu Pro Leu Gly 275 280 285
- Ala Ile Ile Gly Ile Ala Ala Gly Gly Ala Val Leu Leu Leu Leu 290 295 300
- Ala Leu Gly Ile Cys Leu Cys Cys Arg Lys Arg Ser Lys Lys Ala Leu 305 310 315 320
- Gly Asp Pro Glu Ala Thr Thr Ser Ser Arg Arg Pro Trp Phe Thr Pro 325 330 335
- Pro Leu Ser Ala Lys Ser Gln Ser Asp Pro Ser Lys Ser Ile Asp Lys 340 345 350
- Thr Thr Lys Arg Asn Ile Phe Gly Ser Ser Lys Ser Glu Lys Lys Ser 355 360 365
- Ser Lys His Arg Val Phe Glu Pro Ala Pro Leu Asp Lys Gly Ala Ala 370 375 380
- Asp Glu Pro Val Val Lys Ala Ser Pro Pro Val Lys Val Leu Lys Ala 385 390 395 400
- Pro Pro Ser Phe Lys Gly Ile Ser Gly Leu Gly Ala Gly His Ser Lys
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- Ala Thr Ile Gly Lys Val Asn Lys Ser Asn Ile Ala Ala Thr Pro Phe 420 425 430
- Ser Val Ala Asp Leu Gln Ala Ala Thr Asn Ser Phe Ser Gln Asp Asn 435 440 445
- Leu Ile Gly Glu Gly Ser Met Gly Arg Val Tyr Arg Ala Glu Phe Pro 450 455 460
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- Val Gln Asn Glu Asp Asp Phe Leu Ser Val Val Asp Ser Leu Ala Arg 485 490 495
- Leu Gln His Ala Asn Thr Ala Glu Leu Val Gly Tyr Cys Ile Glu His 500 505
- Asp Gln Arg Leu Leu Val Tyr Glu Tyr Val Ser Arg Gly Thr Leu Asn 515 520 525
- Glu Leu Leu His Phe Ser Gly Glu Asn Thr Lys Ala Leu Ser Trp Asn 530 535 540
- Val Arg Ile Lys Ile Ala Leu Gly Ser Ala Arg Ala Leu Glu Tyr Leu 545 550 560
- His Glu Val Cys Ala Pro Pro Val Val His His Asn Phe Lys Ser Ala 570 575
- Asn Ile Leu Leu Asp Asp Glu Leu Asn Pro His Val Ser Asp Cys Gly 580 585
- Leu Ala Ala Leu Ala Pro Ser Gly Ser Glu Arg Gln Val Ser Ala Gln 595 600 605
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- Thr Tyr Thr Val Lys Ser Asp Val Tyr Ser Phe Gly Val Val Met Leu 625 630 635
- Glu Leu Leu Thr Gly Arg Lys Ser Leu Asp Ser Ser Arg Pro Arg Ser 645 650 655
- Glu Gln Ser Leu Val Arg Trp Ala Thr Pro Gln Leu His Asp Ile Asp 660 665
- Ala Leu Ala Arg Met Val Asp Pro Ser Leu Lys Gly Ile Tyr Pro Ala 675 680 685
- Lys Ser Leu Ser Arg Phe Ala Asp Ile Val Ala Leu Cys Val Gln Pro 690 695 700
- Glu Pro Glu Phe Arg Pro Pro Met Ser Glu Val Val Gln Ala Leu Val 705 710 715 720
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- Val Pro Ser Thr Ala Leu Arg Glu Val Ser Leu Leu Gln Met Leu Ser
- His Ser Met Tyr Ile Val Arg Leu Leu Cys Val Glu His Val Glu Lys
- Gly Ser Lys Pro Met Leu Tyr Leu Val Phe Glu Tyr Met Asp Thr Asp
- Leu Lys Lys Tyr Ile Asp Leu His Gly Arg Gly Pro Ser Gly Lys Pro 105
- Leu Pro Pro Lys Val Val Gln Ser Phe Met Tyr Gln Leu Cys Thr Gly
- Leu Ala His Cys His Gly His Gly Val Met His Arg Asp Leu Lys Pro 135
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- Leu Gly Leu Gly Arg Ala Phe Thr Val Pro Met Lys Ser Tyr Thr His 165
- Glu Ile Val Thr Leu Trp Tyr Arg Ala Pro Glu Val Leu Leu Gly Ala 185 180
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<213> Physcomitrella patens

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His Ile Ala Ala Ser Leu Asp Cys Val Pro Val Ala Lys Val Leu Leu

Ala Glu Gly Ala Glu Leu Asn Ala Lys Asp Arg Trp Gly Lys Ser Pro

Arg Gly Glu Ala Glu Ser Ala Gly Tyr Met Glu Met Val Lys Leu Leu

Lys Asp Tyr Gly Ala Glu Ser His Ala Gly Ala Pro Arg Gly His Val 105 1.00

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Lys Gly Ala Phe Gly Glu Ile Arg Lys Ala Leu Trp Arg Gly Thr Pro 150

Val Ala Val Lys Thr Ile Arg Pro Ser Leu Ser Asn Asp Arg Met Val

Ile Lys Asp Phe Gln His Glu Val Gln Leu Leu Val Lys Val Arg His 185

Pro Asn Ile Val Gln Phe Leu Gly Ala Val Thr Arg Gln Arg Pro Leu 200 195

Met Leu Val Thr Glu Phe Leu Ala Gly Gly Asp Leu His Gln Leu Leu 215

Arg Ser Asn Pro Asn Leu Ala Pro Asp Arg Ile Val Lys Tyr Ala Leu 235 225

Asp Ile Ala Arg Gly Met Ser Tyr Leu His Asn Arg Ser Lys Pro Ile 250 245

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Arg Asp Asp Lys Arg Pro Glu Met Arg Ala Gln Thr Tyr Pro Pro Gln

Met Lys Ala Leu Ile Glu Asp Cys Trp Ser Pro Tyr Thr Pro Lys Arg 375

Pro Pro Phe Val Glu Ile Val Lys Lys Leu Glu Val Met Tyr Glu Asp 395

Cys Leu Leu Arg Leu Pro Lys Asp Arg Arg His Leu Arg Asp Ile Leu 410 405

His Leu Arg Arg Asn Pro Ala Asp Ser 420

<210> 30

<211> 283

<212> PRT

<213> Physcomitrella patens

Met Lys Arg Tyr Gln Arg Arg Lys Val Gln Arg Leu Gly Arg Glu Gly

Gln Val Leu Leu Glu Arg Thr Leu Phe Lys Gln Leu Arg Pro Ser Pro 25

Phe Val Pro His Leu Leu Ala Thr Pro Ile Asp Ser Asp Asn Val Ala 40

Leu Val Leu Asn Cys Val Leu Ala Gly Pro Leu Glu Leu Leu Arg

Ser Pro Leu Asp Glu Asn Ser Ala Arg Phe Leu Val Ala Asn Val Val 65

Leu Ala Val Glu Leu Leu His Lys Asp Gly Val Val Tyr Arg Gly Ile

Ser Pro Asp Val Leu Met Ile Asp Arg Lys Gly Arg Leu Gln Leu Val

Asp Phe Arg Phe Ala Lys Gln Met Ser Asp Glu Arg Thr Phe Thr Val

Cys Gly Met Ala Asp Phe Leu Ala Pro Glu Ile Ile Gln Gly Gln Gly 130 135 140

His Gly Leu Ala Ser Asp Trp Trp Ala Val Gly Val Leu Met Tyr Phe 145 150 155 160

Met Leu Gln Thr Glu Leu Pro Phe Gly Ser Trp Arg Asp Asn Glu Leu 165 170 175

Glu Ile Phe Gly Arg Ile Ala Arg Arg Gln Leu Thr Phe Pro Ser Ser 180 185 190

Phe Ser Pro Glu Ala Val Asp Leu Ile Asp Lys Leu Leu Val Val Asp 195 200 205

Pro Thr Lys Arg Leu Gly Cys Asp Ser His Gly Ser Leu Ala Ile Arg 210 215 220

Glu His Pro Trp Phe Arg Gly Ile Asn Trp Asp Lys His Leu Asp Cys 225 230 235 240

Ser Val Glu Val Pro Ser Glu Ile Met Thr Arg Leu Gln Leu Ala Ile 245 250 255

Asp Phe Leu Pro Val Asp Asp Ser Tyr Gln Val Phe Asp Leu Gln Pro 260 265 270

Asp Glu Asp Asp Pro Pro Trp Leu Asp Gly Trp
275
280

<210> 31

<211> 417

<212> PRT

<213> Physcomitrella patens

<400> 31

Met Asp Leu Gly Gly Asp Arg Met Arg Ala Pro Gln Arg Gln Ser Arg
1 5 10 15

Glu Tyr Gln Tyr Arg Ser Leu Asp Val Phe Thr Glu Gln His Glu Gln
20 25 30

Leu Gln Lys Gln Gln Gln Gln Asp Glu Tyr Gln Arg Thr Glu Leu Lys
35 40 45

Leu Glu Thr Leu Pro Lys Met Leu Ser Asn Ala Thr Val Ser Ser Ser 50 55 60

Pro Arg Ser Ser Pro Asp Gly Arg Arg Leu Arg Thr Val Ala Asn Lys 65 70 75 80

- Tyr Ala Val Glu Gly Met Val Gly Ser Gly Ala Phe Cys Lys Val Tyr 85 90 95
- Gln Gly Ser Asp Leu Thr Asn His Glu Val Val Gly Ile Lys Leu Glu 100 105 110
- Asp Thr Arg Thr Glu His Ala Gln Leu Met His Glu Ser Arg Leu Tyr 115 120 125
- Asn Ile Leu Arg Gly Gly Lys Gly Val Pro Asn Met Arg Trp Phe Gly 130 135
- Lys Glu Gln Asp Tyr Asn Val Met Val Leu Asp Leu Leu Gly Pro Asn 145 150 155 160
- Leu Leu His Leu Phe Lys Val Cys Gly Leu Arg Phe Ser Leu Lys Thr 165 170 175
- Val Ile Met Leu Gly Tyr Gln Met Ile Asp Arg Val Glu Tyr Val His 180 185 190
- Ser Arg Gly Leu Val His Arg Asp Leu Lys Pro Asp Asn Phe Leu Met 195 200 205
- Gly Cys Gly Arg Gln Gly Asn Gln Val Phe Ile Ile Asp Phe Gly Leu 210 215 220
- Ala Lys Glu Tyr Met Asp Pro Ala Thr Arg Arg His Ile Pro Tyr Arg 225 230 235 240
- Asp Arg Lys Ser Phe Thr Gly Thr Ala Arg Tyr Ala Ser Arg Asn Gln 245 250
- His Arg Gly Ile Glu His Ser Arg Arg Asp Asp Ile Glu Ser Leu Gly 260 265 270
- Tyr Ile Leu Met Tyr Phe Leu Arg Gly Asn Leu Pro Trp Gln Gly Lys 275 280 285
- Gly Gly Gln Arg Leu Thr Asp Gln Lys Gln His Glu Tyr Met His Asn 290 295 300
- Lys Ile Lys Met Asn Thr Thr Val Glu Glu Leu Cys Asp Gly Tyr Pro 305 310 310
- Ser Gln Phe Ala Asp Phe Leu His His Ala Arg Ser Leu Gly Phe Tyr 325 330 335
- Glu Gln Pro Asp Tyr Cys Tyr Leu Arg Ser Leu Phe Arg Asp Leu Phe 340
- Ile Gln Lys Lys Phe Gln Leu Asp His Val Tyr Asp Trp Thr Val Tyr 355 360 365
- Thr Gln Leu Pro Gln Asn Gly Ser Leu Gln Ser Val Arg Ser Gln Asn 370 375 380

Ser Ala Ala Ser Ser His Leu Gln Asn Arg Pro Ser Asn Val Ser Tyr 385 390 395 400

Cys Pro Pro Leu Thr Lys Ser Glu Phe Arg Arg Glu Val Val Ala Ala 405 410 415

Asn

<210> 32

<211> 484

<212> PRT

<213> Physcomitrella patens

<400> 32

Met Glu Pro Arg Val Gly Asn Lys Tyr Arg Leu Gly Arg Lys Ile Gly
1 5 10 15

Ser Gly Ser Phe Gly Glu Ile Tyr Leu Gly Thr Asn Val Gln Thr Asn 20 25 30

Glu Glu Val Gly Ile Lys Leu Glu Ser Ile Lys Thr Lys His Pro Gln
35 40 45

Leu Leu Tyr Glu Ser Lys Leu Tyr Arg Ile Leu Gln Gly Gly Thr Gly
50 55 60

Ile Pro Asn Ile Arg Trp Phe Gly Ile Glu Gly Asp Tyr Asn Val Leu 65 70 75 80

Val Leu Asp Leu Gly Pro Ser Leu Glu Asp Leu Phe Asn Phe Cys
85 90 95

Ser Arg Lys Phe Ser Leu Lys Thr Val Leu Met Leu Ala Asp Gln Leu 100 105 110

Ile Asn Arg Val Glu Tyr Val His Ala Lys Ser Phe Leu His Arg Asp 115 120 125

Ile Lys Pro Asp Asn Phe Leu Met Gly Leu Gly Arg Arg Ala Asn Gln 130 135 140

Val Tyr Ile Ile Asp Phe Gly Leu Ala Lys Lys Tyr Arg Asp Pro Ser 145 150 155 160

Thr His Gln His Ile Pro Tyr Arg Glu Asn Lys Asn Leu Thr Gly Thr
165 170 175

Ala Arg Tyr Ala Ser Ile Asn Thr His Leu Gly Ile Glu Gln Ser Arg 180 185 190

Arg Asp Asp Leu Glu Ser Leu Gly Tyr Val Leu Met Tyr Phe Leu Arg 195 200 205

Gly Ser Leu Pro Trp Gln Gly Leu Lys Ala Gly Thr Lys Lys Gln Lys 210 215 220

Tyr Glu Lys Ile Ser Glu Lys Lys Met Ser Thr Pro Ile Glu Val Leu 225 230 235 240

Cys Lys Asn Tyr Pro Ser Glu Phe Ala Ser Tyr Phe His Tyr Cys Arg 245 250 255

Ser Leu Arg Phe Asp Asp Lys Pro Asp Tyr Ala Tyr Leu Lys Arg Ile 260 265 270

Phe Arg Asp Leu Phe Ile Arg Glu Gly Phe Gln Phe Asp Tyr Val Phe 275 280 285

Asp Trp Thr Ile Leu Lys Tyr Gln Gln Ser Gln Ile Ser Gly Gly Ser 290 295 300

Ser Thr Arg Leu Gly Ala Ser Ala Gly Gln Thr Ser Gly Ala Leu Gly 305 310 315 320

Thr Gly Ala Thr Gly Ser Arg Asp Leu Gln Arg Pro Thr Glu Pro Met 325 330 335

Asp Pro Ser Arg Arg Leu Pro Gly Gly Ala Asn Gly Ser Gly Val 340 345

Ala Asn Ala Leu Asp Ser Ser Lys His Lys Ser Pro Gly Leu Asp Glu 355

Ser Ala Lys Asp Ser Ala Leu Ala Val Val Ser Glu Pro Glu Arg Met 370 375 380

His Thr Ser Ser Tyr Ala Thr Arg Gly Gly Ser Ser Ser Arg Arg Ala 385 390 395 400

Val Leu Ser Ser Ser Arg Pro Ser Gly Ala Ser Ala Glu Val Val Asp 405 410 415

Ser Ser Arg Thr Gly Ser Ser Lys Leu Gly Pro Thr Ser Leu Arg Ser 420 425 430

Ser Ala Gly Met Gln Arg Ser Ser Pro Val Thr Ser Asp Pro Lys Arg 435 440 445

Ile Ser Ser Arg His Pro Gln Pro Pro Ser Ala Asn Leu Arg Ile Tyr 450 455 460

Glu Ala Ala Ile Lys Gly Val Glu Ser Leu Ser Val Glu Val Asp Gln 465 470 475 480

Ser Arg Tyr Lys

<210> 33

<211> 333

<212> PRT

<213> Physcomitrella patens

- Met Ser Lys Ala Arg Val Tyr Thr Asp Val Asn Val Gln Arg Pro Lys
- Asp Tyr Trp Asp Tyr Glu Ala Leu Thr Val Gln Trp Gly Asp Gln Asp
- Asp Tyr Glu Val Val Arg Lys Val Gly Arg Gly Lys Tyr Ser Glu Val
- Phe Glu Gly Val Asn Ala Val Asn Ser Glu Arg Cys Val Met Lys Ile 55
- Leu Lys Pro Val Lys Lys Lys Ile Lys Arg Glu Ile Lys Ile Leu
- Gln Asn Leu Cys Gly Gly Pro Asn Ile Val Lys Leu Leu Asp Ile Val
- Arg Asp Gln Gln Ser Lys Thr Pro Ser Leu Ile Phe Glu Tyr Val Asn 105
- Asn Thr Asp Phe Lys Val Leu Tyr Pro Thr Leu Thr Asp Phe Asp Ile 120 115
- Arg Tyr Tyr Ile His Glu Leu Leu Lys Ala Leu Asp Tyr Cys His Ser 135
- Gln Gly Ile Met His Arg Asp Val Lys Pro His Asn Val Met Ile Asp 150
- His Glu Gln Arg Lys Leu Arg Leu Ile Asp Trp Gly Leu Ala Glu Phe
- Tyr His Pro Gly Lys Glu Tyr Asn Val Arg Val Ala Ser Arg Tyr Phe
- Lys Gly Pro Glu Leu Leu Val Asp Leu Gln Asp Tyr Asp Tyr Ser Leu 200
- Asp Met Trp Ser Leu Gly Cys Met Phe Ala Gly Met Ile Phe Arg Lys
- Glu Pro Phe Phe Tyr Gly His Asp Asn Tyr Asp Gln Leu Val Lys Ile 230
- Ala Lys Val Leu Gly Thr Asp Glu Leu Asn Ser Tyr Leu Asn Lys Tyr
- Arg Leu Glu Leu Asp Pro His Leu Glu Ala Leu Val Gly Arg His Ser 265
- Arg Lys Pro Trp Ser Lys Phe Ile Asn Ala Asp Asn Gln Arg Leu Val 280 275
- Val Pro Glu Ala Val Asp Phe Leu Asp Lys Leu Leu Arg Tyr Asp His 300 295 290

Gln Asp Arg Leu Thr Ala Lys Glu Ala Met Ala His Pro Tyr Phe Tyr 305 310 315 320

Pro Val Lys Val Ser Glu Val Ser Asn Arg Arg Ser Ala 325 330

<210> 34

<211> 375

<212> PRT

<213> Physcomitrella patens

<400> 34

Met Glu Thr Ser Ser Gly Thr Pro Glu Leu Lys Val Ile Ser Thr Pro 1 5 10 15

Thr Tyr Gly Gly His Tyr Val Lys Tyr Val Val Ala Gly Thr Asp Phe 20 25 30

Glu Val Thr Ala Arg Tyr Lys Pro Pro Leu Arg Pro Ile Gly Arg Gly
35 40 45

Ala Tyr Gly Ile Val Cys Ser Leu Phe Asp Thr Val Thr Gly Glu Glu 50 55 60

Val Ala Val Lys Lys Ile Gly Asn Ala Phe Asp Asn Arg Ile Asp Ala 65 70 75 80

Lys Arg Thr Leu Arg Glu Ile Lys Leu Leu Arg His Met Asp His Glu 85 90 95

Asn Val Val Ala Ile Thr Asp Ile Ile Arg Pro Pro Thr Arg Glu Asn 100 105 110

Phe Asn Asp Val Tyr Ile Val Tyr Glu Leu Met Asp Thr Asp Leu His
115 120 125

Gln Ile Ile Arg Ser Asn Gln Ala Leu Thr Glu Asp His Cys Gln Tyr 130 135 140

Phe Leu Tyr Gln Ile Leu Arg Gly Leu Lys Tyr Ile His Ser Ala Asn 145 150 155 160

Val Leu His Arg Asp Leu Lys Pro Thr Asn Leu Leu Val Asn Ala Asn 165 170 175

Cys Asp Leu Lys Ile Ala Asp Phe Gly Leu Ala Arg Thr Leu Ser Glu 180 185 190

Thr Asp Phe Met Thr Glu Tyr Val Val Thr Arg Trp Tyr Arg Ala Pro 195 200 205

Glu Leu Leu Asn Cys Ser Ala Tyr Thr Ala Ala Ile Asp Ile Trp 210 215 220

Ser Val Gly Cys Ile Phe Met Glu Leu Leu Asn Arg Ser Ala Leu Phe 225 230 235 240 Pro Gly Arg Asp Tyr Val His Gln Leu Arg Leu Ile Thr Glu Leu Ile 245 250 255

Gly Thr Pro Glu Asp Arg Asp Leu Gly Phe Leu Arg Ser Asp Asn Ala 260 265 270

Arg Arg Tyr Ile Lys His Leu Pro Arg Gln Ser Pro Ile Pro Leu Thr 275 280 285

Gln Lys Phe Arg Gly Ile Asn Arg Ser Ala Leu Asp Leu Val Glu Lys 290 295 300

Met Leu Val Phe Asp Pro Ala Lys Arg Ile Thr Val Glu Ala Ala Leu 305 310 315 320

Ala His Pro Tyr Leu Ala Ser Leu His Asp Ile Asn Asp Glu Pro Ala 325 330 335

Ser Val Ser Pro Phe Glu Phe Asp Phe Glu Glu Pro Pro Ile Ser Glu 340 345 350

Glu His Ile Lys Asp Leu Ile Trp Arg Glu Ala Leu Asp Cys Ser Leu 355 360 365

Gly Pro Asp Asp Met Val Gln 370 375

<210> 35

<211> 331

<212> PRT

<213> Physcomitrella patens

<400> 35

Met Gly Leu Thr Pro Phe Ser Cys Val Thr Val Gln Gly Tyr Val Arg

1 5 10 15

Val Val Tyr Pro Asp Gly His Val Glu Asn Leu Ser Lys Ser Cys Ser 20 25 30

Val His Asp Leu Leu Gly Asn Pro Asp Tyr Tyr Val Cys Gly Ser

Thr Pro Tyr Thr Ile Thr Asn Arg Met Ala Ala Glu Glu Val Leu Glu 50 55 60

Tyr Gly Val Thr Tyr Phe Val Cys Ala Thr Pro Asn Ala Gln Pro Phe 65 70 75 80

Leu Glu Arg Gln Pro Lys Val Val His Arg Gly Ser Lys Ile Leu Pro 85 90 95

Arg Phe Ser Lys His Gly Val His Val Arg Glu Leu Arg Ser Pro Thr
100 105 110

His Gly Ser Gln Gln Ser Arg Lys Val Phe Asp Tyr His Ser Val Thr 115 120 125 Met Gln Gln Leu Glu Ser Ile Arg Asn Glu Gly Pro Glu Pro His Leu 130 135 140

Ala Gly Asp Arg Pro Ser Lys His Leu Lys Leu Val Phe Ile Arg His 145 150 155 160

Cys Leu Arg Ala Leu Arg Leu Pro Arg Ile Ser Ile Asp Leu Met Glu 165 170 175

Ser Pro Leu Pro Asn Leu Ser Gly Glu Ala Leu Ser Pro Thr Ala Thr 180 185 190

Ala Lys Asp Glu Ile Thr Gln Met Ile Leu Lys Ser Ala Ala Arg Ser 195 200 205

Glu Leu Gly Met Tyr Val Ser Lys Arg Gln Glu Phe Tyr Leu Arg Arg 210 215 220

Ala Arg Arg Arg Lys Phe Ala Trp Lys Pro Val Leu Gln Ser Ile 225 230 235 240

Ser Glu Met Lys Pro Val Met Glu Phe His Thr Pro Met Ala Tyr Arg 245 250 255

Asp Ser Gly Ser Pro Pro Lys Asn Ala Ser Thr Pro Ser Leu Pro Gly 260 265 270

Pro Lys Asn Ile Ser Pro Pro Arg Gln Val Ser Val Pro Gln Arg Ser 275 280 285

Ser Pro Pro Pro Lys Asn Val Ser Pro Pro Pro Gln Pro Ala Phe Val 290 295 300

Ala Arg Thr Ala Ser Lys Tyr Ser Ala Ala Ser Gln Gln Val Gln Arg 305 310 315 320

Asn Arg Gly Asn Ala Lys Ser Leu Tyr Met Ala 325 330

<210> 36

<211> 346

<212> PRT

<213> Physcomitrella patens

<400> 36

Met Ser Arg Arg Val Arg Arg Gly Gly Leu Arg Val Ala Val Pro Lys
1 5 10 15

Gln Glu Thr Pro Val Ser Lys Phe Leu Thr Ala Ser Gly Thr Phe Gln
20 25 30

Asp Asp Asp Ile Lys Leu Asn His Thr Gly Leu Arg Val Val Ser Ser 35 40 45

Glu Pro Asn Leu Pro Thr Gln Thr Gln Ser Ser Pro Asp Gly Gln
50 60

	Ser	Ile	Ala	Asp	Leu 70	Glu	Leu	Val	Arg	Phe 75	Leu	Gly	Lys	Gly	Ala 80
65	501	110		E	70					75					8

- Gly Gly Thr Val Gln Leu Val Arg His Lys Trp Thr Asn Val Asn Tyr 85 90 95
- Ala Leu Lys Ala Ile Gln Met Asn Ile Asn Glu Thr Val Arg Lys Gln 100 105 110
- Ile Val Gln Glu Leu Lys Ile Asn Gln Val Thr His Gln Gln Cys Pro 115 120 125
- Tyr Ile Val Glu Cys Phe His Ser Phe Tyr His Asn Gly Val Ile Ser 130 135 140
- Met Ile Leu Glu Tyr Met Asp Arg Gly Ser Leu Ser Asp Ile Ile Lys 145 150 155 160
- Gln Gln Lys Gln Ile Pro Glu Pro Tyr Leu Ala Val Ile Ala Ser Gln 165 170 175
- Val Leu Lys Gly Leu Glu Tyr Leu His Gln Val Arg His Ile Ile His
 180 185 190
- Arg Asp Ile Lys Pro Ser Asn Leu Leu Ile Asn His Lys Gly Glu Val 195 200 205
- Lys Ile Ser Asp Phe Gly Val Ser Ala Val Leu Val His Ser Leu Ala 210 215 220
- Gln Arg Asp Thr Phe Val Gly Thr Cys Thr Tyr Met Ser Pro Glu Arg 225 230 235
- Leu Gln Gly Arg Ser Tyr Ala Tyr Asp Ser Asp Leu Trp Ser Leu Gly 245 250 255
- Leu Thr Leu Leu Glu Cys Ala Leu Gly Thr Phe Pro Tyr Lys Pro Ala 260 265 270
- Gly Met Glu Glu Gly Trp Gln Asn Phe Phe Ile Leu Met Glu Cys Ile 275 280 285
- Val Asn Gln Pro Pro Ala Ala Ala Ser Pro Asp Lys Phe Ser Pro Glu 290 295 300
- Phe Cys Ser Phe Ile Glu Ser Cys Ile Arg Lys Cys Pro Ser Glu Arg 305 310 315
- Pro Ser Thr Thr Asp Leu Leu Lys His Pro Phe Leu Gln Lys Tyr Asn 325 330 335
- Glu Glu Glu Tyr His Leu Ser Lys Ile Leu 340 345

<210> 37

<211> 346

<212> PRT

<213> Physcomitrella patens

<400> 37

Met Ser Arg Arg Val Arg Arg Gly Gly Leu Arg Val Ala Val Pro Lys

Gln Glu Thr Pro Val Ser Lys Phe Leu Thr Ala Ser Gly Thr Phe Gln 25

Asp Asp Asp Ile Lys Leu Asn His Thr Gly Leu Arg Val Val Ser Ser

Glu Pro Asn Leu Pro Thr Gln Thr Gln Ser Ser Pro Asp Gly Gln

Leu Ser Ile Ala Asp Leu Glu Leu Val Arg Phe Leu Gly Lys Gly Ala 65

Gly Gly Thr Val Gln Leu Val Arg His Lys Trp Thr Asn Val Asn Tyr

Ala Leu Lys Ala Ile Gln Met Asn Ile Asn Glu Thr Val Arg Lys Gln 105

Ile Val Gln Glu Leu Lys Ile Asn Gln Val Thr His Gln Gln Cys Pro 120

Tyr Ile Val Glu Cys Phe His Ser Phe Tyr His Asn Gly Val Ile Ser 135 130

Met Ile Leu Glu Tyr Met Asp Arg Gly Ser Leu Ser Asp Ile Ile Lys

Gln Gln Lys Gln Ile Pro Glu Pro Tyr Leu Ala Val Ile Ala Ser Gln 165

Val Leu Lys Gly Leu Glu Tyr Leu His Gln Val Arg His Ile Ile His 185

Arg Asp Ile Lys Pro Ser Asn Leu Leu Ile Asn His Lys Gly Glu Val

Lys Ile Ser Asp Phe Gly Val Ser Ala Val Leu Val His Ser Leu Ala

Gln Arg Asp Thr Phe Val Gly Thr Cys Thr Tyr Met Ser Pro Glu Arg 235

Leu Gln Gly Arg Ser Tyr Ala Tyr Asp Ser Asp Leu Trp Ser Leu Gly

Leu Thr Leu Leu Glu Cys Ala Leu Gly Thr Phe Pro Tyr Lys Pro Ala 265

Gly Met Glu Glu Gly Trp Gln Asn Phe Phe Ile Leu Met Glu Cys Ile 280 275

Val Asn Gln Pro Pro Ala Ala Ala Ser Pro Asp Lys Phe Ser Pro Glu 290 295 300

Phe Cys Ser Phe Ile Glu Ser Cys Ile Arg Lys Cys Pro Ser Glu Arg 305 310 315 320

Pro Ser Thr Thr Asp Leu Lys His Pro Phe Leu Gln Lys Tyr Asn 325 330 335

Glu Glu Glu Tyr His Leu Ser Lys Ile Leu 340 345

<210> 38

<211> 597

<212> PRT

<213> Physcomitrella patens

<400> 38

Met Gly Gln Cys Tyr Gly Lys Phe Asp Asp Gly Glu Gly Glu Asp
1 5 10 15

Leu Phe Glu Arg Gln Lys Val Gln Val Ser Arg Thr Pro Lys His Gly
20 25 30

Ser Trp Ser Asn Ser Asn Arg Gly Ser Phe Asn Asn Gly Gly Ala
35 40 45

Ser Pro Met Arg Ala Lys Thr Ser Phe Gly Ser Ser His Pro Ser Pro 50 55 60

Arg His Pro Ser Ala Ser Pro Leu Pro His Tyr Thr Ser Ser Pro Ala 65 70 75 80

Pro Ser Thr Pro Arg Arg Asn Ile Phe Lys Arg Pro Phe Pro Pro 85 90 95

Ser Pro Ala Lys His Ile Gln Ser Ser Leu Val Lys Arg His Gly Ala 100 105 110

Lys Pro Lys Glu Gly Gly Ala Ile Pro Glu Ala Val Asp Gly Glu Lys 115 120 125

Pro Leu Asp Lys His Phe Gly Tyr His Lys Asn Phe Ala Thr Lys Tyr 130 135 140

Glu Leu Gly His Glu Val Gly Arg Gly His Phe Gly His Thr Cys Tyr 145 150 155 160

Ala Lys Val Arg Lys Gly Glu His Lys Gly Gln Ala Val Ala Val Lys 165 170 175

Ile Ile Ser Lys Ala Lys Met Thr Thr Ala Ile Ala Ile Glu Asp Val

Gly Arg Glu Val Lys Ile Leu Lys Ala Leu Thr Gly His Gln Asn Leu 195 200 205

- Val Arg Phe Tyr Asp Ser Cys Glu Asp His Leu Asn Val Tyr Ile Val 210 215 220
- Met Glu Leu Cys Glu Gly Gly Glu Leu Leu Asp Arg Ile Leu Ser Arg 225 230 235 240
- Gly Gly Lys Tyr Ser Glu Glu Asp Ala Lys Val Val Val Arg Gln Ile 245 250 255
- Leu Ser Val Val Ala Phe Cys His Leu Gln Gly Val Val His Arg Asp 260 265 270
- Leu Lys Pro Glu Asn Phe Leu Phe Thr Thr Lys Asp Glu Tyr Ala Gln 275 280 285
- Leu Lys Ala Ile Asp Phe Gly Leu Ser Asp Phe Ile Lys Pro Asp Glu 290 295 300
- Arg Leu Asn Asp Ile Val Gly Ser Ala Tyr Tyr Val Ala Pro Glu Val 305 310 315
- Leu His Arg Leu Tyr Ser Met Glu Ala Asp Val Trp Ser Ile Gly Val 325
- Ile Thr Tyr Ile Leu Leu Cys Gly Ser Arg Pro Phe Trp Ala Arg Thr 340 345 350
- Glu Ser Gly Ile Phe Arg Ala Val Leu Arg Ala Asp Pro Ser Phe Glu 355 360 365
- Glu Ala Pro Trp Pro Ser Ile Ser Pro Glu Ala Lys Asp Phe Val Lys 370 375 380
- Arg Leu Leu Asn Lys Asp Met Arg Lys Arg Met Thr Ala Ala Gln Ala 385 390 395 400
- Leu Thr His Pro Trp Ile Arg Ser Asn Asn Val Lys Ile Pro Leu Asp 405 410 415
- Ile Leu Val Tyr Arg Leu Val Arg Asn Tyr Leu Arg Ala Ser Ser Met 420 425 430
- Arg Lys Ala Ala Leu Lys Ala Leu Ser Lys Thr Leu Thr Glu Asp Glu
 435 440 445
- Thr Phe Tyr Leu Arg Thr Gln Phe Met Leu Leu Glu Pro Ser Asn Asn 450 455 460
- Gly Arg Val Thr Phe Glu Asn Phe Arg Gln Ala Leu Leu Lys Asn Ser 475 480
- Thr Glu Ala Met Lys Glu Ser Arg Val Phe Glu Ile Leu Glu Ser Met 485 490 495
- Asp Gly Leu His Phe Lys Lys Met Asp Phe Ser Glu Phe Cys Ala Ala 500 505 510

Ala Ile Ser Val Leu Gln Leu Glu Ala Thr Glu Arg Trp Glu Gln His 520

Ala Arg Ala Ala Tyr Asp Ile Phe Glu Lys Glu Gly Asn Arg Val Ile

Tyr Pro Asp Glu Leu Ala Lys Glu Met Gly Leu Ala Pro Asn Val Pro

Ala Gln Val Phe Leu Asp Trp Ile Arg Gln Ser Asp Gly Arg Leu Ser 570 565

Phe Thr Gly Phe Thr Lys Leu Leu His Gly Ile Ser Ser Arg Ala Ile 585

Lys Asn Leu Gln Gln 595

<210> 39

<211> 549

<212> PRT

<213> Physcomitrella patens

<400> 39

Met Gly Asn Thr Ser Ser Arg Gly Ser Arg Lys Ser Thr Arg Gln Val 10

Asn Gln Gly Val Gly Ser Gln Asp Thr Arg Glu Lys Asn Asp Ser Val

Asn Pro Lys Thr Arg Gln Gly Gly Ser Val Gly Ala Asn Asn Tyr Gly

Gly Lys Pro Ser Ser Gly Ala Gln Ala Gly Glu Arg Ser Thr Ser Ala

Pro Ala Ala Leu Pro Arg Pro Lys Pro Ala Ser Arg Ser Val Ser Gly 70

Val Leu Gly Lys Pro Leu Ser Asp Ile Arg Gln Ser Tyr Ile Leu Gly

Arg Glu Leu Gly Arg Gly Gln Phe Gly Val Thr Tyr Leu Cys Thr Asp 105

Lys Met Thr Asn Glu Ala Tyr Ala Cys Lys Ser Ile Ala Lys Arg Lys 120

Leu Thr Ser Lys Glu Asp Ile Glu Asp Val Lys Arg Glu Val Gln Ile 135

Met His His Leu Ser Gly Thr Pro Asn Ile Val Val Leu Lys Asp Val 155 145

Phe Glu Asp Lys His Ser Val His Leu Val Met Glu Leu Cys Ala Gly 170 165

- Gly Glu Leu Phe Asp Arg Ile Ile Ala Lys Gly His Tyr Ser Glu Arg 180 185 190
- Ala Ala Asp Met Cys Arg Val Ile Val Asn Val Val His Arg Cys 195 200 205
- His Ser Leu Gly Val Phe His Arg Asp Leu Lys Pro Glu Asn Phe Leu 210 215 220
- Leu Ala Ser Lys Ala Glu Asp Ala Pro Leu Lys Ala Thr Asp Phe Gly 225 230 235 240
- Leu Ser Thr Phe Phe Lys Pro Gly Asp Val Phe Gln Asp Ile Val Gly 245 250 255
- Ser Ala Tyr Tyr Val Ala Pro Glu Val Leu Lys Arg Ser Tyr Gly Pro 260 265 270
- Glu Ala Asp Val Trp Ser Ala Gly Val Ile Val Tyr Ile Leu Leu Cys 275 280 285
- Gly Val Pro Pro Phe Trp Ala Glu Thr Glu Gln Gly Ile Phe Asp Ala 290 295 300
- Val Leu Lys Gly His Ile Asp Phe Glu Asn Asp Pro Trp Pro Lys Ile 305 310 315 320
- Ser Asn Gly Ala Lys Asp Leu Val Arg Lys Met Leu Asn Pro Asn Val 325 330 335
- Lys Ile Arg Leu Thr Ala Gln Gln Val Leu Asn His Pro Trp Met Lys 340 345 350
- Glu Asp Gly Asp Ala Pro Asp Val Pro Leu Asp Asn Ala Val Leu Thr 355 360 365
- Arg Leu Lys Asn Phe Ser Ala Ala Asn Lys Met Lys Lys Leu Ala Leu 370 375 380
- Lys Val Ile Ala Glu Ser Leu Ser Glu Glu Glu Ile Val Gly Leu Arg 385 390 395 400
- Glu Met Phe Lys Ser Ile Asp Thr Asp Asn Ser Gly Thr Val Thr Phe 405 410 415
- Glu Glu Leu Lys Glu Gly Leu Leu Lys Gln Gly Ser Lys Leu Asn Glu
 420 425 430
- Ser Asp Ile Arg Lys Leu Met Glu Ala Ala Asp Val Asp Gly Asn Gly 435 440 445
- Lys Ile Asp Phe Asn Glu Phe Ile Ser Ala Thr Met His Met Asn Lys 450 450
- Thr Glu Lys Glu Asp His Leu Trp Ala Ala Phe Met His Phe Asp Thr 465 470 475 480

Asp Asn Ser Gly Tyr Ile Thr Ile Asp Glu Leu Gln Glu Ala Met Glu 490 Lys Asn Gly Met Gly Asp Pro Glu Thr Ile Gln Glu Ile Ile Ser Glu 505 Val Asp Thr Asp Asn Asp Gly Arg Ile Asp Tyr Asp Glu Phe Val Ala 520 Met Met Arg Lys Gly Asn Pro Gly Ala Glu Asn Gly Gly Thr Val Asn Lys Pro Arg His Arg 545 <210> 40 <211> 18 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Primer <400> 40 18 caggaaacag ctatgacc <210> 41 <211> 19 <212> DNA <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Primer <400> 41 19 ctaaagggaa caaaagctg <210> 42 <211> 18 <212> DNA <213> Artificial Sequence <223> Description of Artificial Sequence: Primer <400> 42 tgtaaaacga cggccagt 18 <210> 43 <211> 25 <212> DNA <213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Primer	
<400> 43	
ccacggtctt cggctgctgg tcgtg	25
<210> 44	
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gcagcacagc accaccagcg gctat	23
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<210> 47 <211> 33	
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gcgttaactc gaccaaggtc actattccaa gca	33
-	
<210> 48	
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sprox profit ordiner	

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<223> Description of Artificial Sequence: Primer
<400> 48
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cggtgcccac ctcgttcctg tggtt
<210> 49
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<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 49
                                                                   31
atcccgggag tgggtggttg gactgtaagg a
<210> 50
<211> 34
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 50
gcgttaacct tcgtcttgga caggtagagg ttac
                                                                   34
<210> 51
<211> 25
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 51
                                                                   25
gactcagccc cgtaatcctt caaca
<210> 52
<211> 31
<212> DNA
<213> Artificial Sequence
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<400> 52
atcccgggca acgagaagca ttcgagatgg c
                                                                   31
<210> 53
<211> 33
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<223> Description of Artificial Sequence: Primer	
<400> 53	
gcgttaacga gcatcacgat actcggtgat ttc	33
-010. 54	
<210> 54 <211> 27	
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(223) bescription of Artificial Sequence. Filmer	
<400> 54 cgacggctaa taccacgttg gegacca	27
<210> 55	
<211> 33	
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<210> 56	
<211> 34 <212> DNA	
<212> DNA <213> Artificial Sequence	
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gcgagctcgc accactgaat gatggagact cagg	34
<210> 57	
<211> 25 <212> DNA	
<213> Artificial Sequence	
<220>	
<223> Description of Artificial Sequence: Primer	
<400> 57	
cgaccgcagc ccatgaggaa gttat	25

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<210> 58
<211> 33
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Primer
<400> 58
atcccgggct cacgtagtgc actgaactct gtc
                                                                    33
<210> 59
<211> 33
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Primer
<400> 59
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gcgttaacat gcccatcttc tcatactcag acc
<210> 60
<211> 25
<212> DNA
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<223> Description of Artificial Sequence: Primer
<400> 60
ctcgcctacc aagccccatt agaaa
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<210> 61
<211> 32
<212> DNA
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<223> Description of Artificial Sequence: Primer
<400> 61
atcccgggtt gtcgaggacg gagagagaag ag
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<210> 62
<211> 33
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<223> Description of Artificial Sequence: Primer
<400> 62
gcgttaacct taggaatcgt atggcagaga gct
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<210> 63
<211> 25
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 63
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gcttcacaat gttgggccct ccaca
<210> 64
<211> 33
<212> DNA
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<223> Description of Artificial Sequence: Primer
<400> 64
gcgttaacgg gaggaaggtc gggggaagag acg
                                                                    33
<210> 65
<211> 33
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<223> Description of Artificial Sequence: Primer
<400> 65
gcgagctcag cgcttcgcac aactgagaaa cct
                                                                    33
<210> 66
<211> 25
<212> DNA
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<223> Description of Artificial Sequence: Primer
<400> 66
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acgagaaggt tggtgggctt caagt
<210> 67
<211> 30
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Primer
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atcccgggcg agccatggcg ccacttgctt
<210> 68
<211> 33
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<220>
<223> Description of Artificial Sequence: Primer
<400> 68
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gcgttaacgc cgagcaacaa tgtctgctgg atg
<210> 69
<211> 25
<212> DNA
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<223> Description of Artificial Sequence: Primer
<400> 69
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cccggtaagc catcggagtg tggaa
<210> 70
<211> 30
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 70
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atcccgggct tgtattggct cggataattt
<210> 71
<211> 33
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<223> Description of Artificial Sequence: Primer
<400> 71
gcgttaacgg caatatctgc acagccgttc act
                                                                    33
<210> 72
<211> 25
<212> DNA
<213> Artificial Sequence
<220>
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<223> Description of Artificial Sequence: Primer
<400> 72
                                                                    25
gtgtctcgct gggccaagga atgaa
<210> 73
<211> 35
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Primer
<400> 73
                                                                    35
atcccgggcg gtcgagtcgt attaggtgtt gtttc
<210> 74
<211> 30
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 74
gageteeggt aggteegace tetteaattg
                                                                    30
<210> 75
<211> 26
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 75
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gacgacgcga agcccggtgt ggttga
<210> 76
<211> 31
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 76
                                                                    31
atcccgggag aggctgatct gatgctacag t
<210> 77
<211> 33
<212> DNA
<213> Artificial Sequence
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<220> <223> Description of Artificial Sequence: Primer	
<400> 77 atgagetetg geggattgge gaggtagtte gae	33
<210> 78 <211> 25 <212> DNA	
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<223> Description of Artificial Sequence: Primer	
<400> 78 cggcgcaacg tagtatgcgc ttcca	25
<210> 79	
<211> 27 <212> DNA	
<213> Artificial Sequence	
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<400> 79	27
cgcggtgaac aacaccttgc aggtgac	27
<210> 80 <211> 25	
<212> DNA <213> Artificial Sequence	
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<400> 80 gctcgggtca gccctcaaca ccgca	25
<210> 81 <211> 25 <212> DNA	
<213> Artificial Sequence	
<220> <223> Description of Artificial Sequence: Primer	
<400> 81 gttaaagctt gtgcagcagt catgc	25
<210> 82 <211> 31	

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<213> Artificial Sequence
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<223> Description of Artificial Sequence: Primer
<400> 82
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atcccgggtg taggcgggcg aggttcgatg c
<210> 83
<211> 34
<212> DNA
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<223> Description of Artificial Sequence: Primer
<400> 83
gcgttaacga caaccggagt agaacggcag tcca
                                                                    34
<210> 84
<211> 25
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 84
agaagcgagg aatgggcagg gacga
                                                                    25
<210> 85
<211> 32
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<223> Description of Artificial Sequence: Primer
<400> 85
atcccgggcg aactgcgatc tgagattcca ac
                                                                    32
<210> 86
<211> 34
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 86
gcgttaacga gatccaaccg aagccatcct acga
                                                                    34
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<210> 87
<211> 30
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Primer
<400> 87
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gcgctgcaga tttcatttgg agaggacacg
<210> 88
<211> 35
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Primer
<400> 88
                                                                    35
cgcggccggc ctcagaagaa ctcgtcaaga aggcg
<210> 89
<211> 25
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 89
                                                                    25
gctgacacgc caagectege tagte
<210> 90
<211> 33
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 90
                                                                    33
gcgttaactc gaccaaggtc actattccaa gca
<210> 91
<211> 34
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 91
                                                                    34
gcgttaacct tcgtcttgga caggtagagg ttac
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<210> 92
<211> 33
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 92
                                                                    33
gcgttaacga gcatcacgat actcggtgat ttc
<210> 93
<211> 34
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 93
                                                                    34
gcgagctcgc accactgaat gatggagact cagg
<210> 94
<211> 33
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 94
                                                                    33
gcgttaacat gcccatcttc tcatactcag acc
<210> 95
<211> 33
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Primer
<400> 95
                                                                    33
gcgttaacct taggaatcgt atggcagaga gct
<210> 96
<211> 33
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
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<400> 96
gcgagctcag cgcttcgcac aactgagaaa cct
                                                                    33
<210> 97
<211> 33
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 97
                                                                    33
gcgttaacgg caatatctgc acagccgttc act
<210> 98
<211> 33
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 98
                                                                    33
gcgttaacgg caatatctgc acagccgttc act
<210> 99
<211> 30
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 99
                                                                    30
gageteeggt aggteegace tetteaattg
<210> 100
<211> 33
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 100
atgagetetg geggattgge gaggtagtte gae
                                                                    33
<210> 101
<211> 34
<212> DNA
<213> Artificial Sequence
<220>
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<223> Description of Artificial Sequence: Primer
<400> 101
gcgttaacga caaccggagt agaacggcag tcca
                                                                    34
<210> 102
<211> 34
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 102
gcgttaacga gatccaaccg aagccatcct acga
                                                                    34
<210> 103
<211> 25
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 103
                                                                    25
cccagtaata gcagggttgg aggaa
<210> 104
<211> 25
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 104
                                                                    25
ggctgcctga agatccgcta cagag
<210> 105
<211> 25
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 105
                                                                    25
cgtcaggcta ctttgcgtgg agcac
<210> 106
<211> 25
<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Primer
<400> 106
cggtgctggc taacaccagg ccaga
                                                                    25
<210> 107
<211> 31
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 107
                                                                    31
atcccgggca acgagaagca ttcgagatgg c
<210> 108
<211> 33
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Primer
<400> 108
                                                                    33
gcgttaacga gcatcacgat actcggtgat ttc
<210> 109
<211> 25
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 109
                                                                    25
cgtggcatct ctcccgatgt tctta
<210> 110
<211> 25
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 110
ggccaactga aggcgtgtca tgatc
                                                                    25
<210> 111
<211> 25
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<212> DNA <213> Artificial Sequence	
<220> <223> Description of Artificial Sequence: Primer	
<400> 111 ctcgagggct cgttcaccgt gacct	25
<210> 112 <211> 26	
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<220> <223> Description of Artificial Sequence: Primer	
<400> 112 cggaggtaac agtagtcagg ctgctc	26
<210> 113 <211> 25 <212> DNA	
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<220> <223> Description of Artificial Sequence: Primer	
<400> 113 ccgcgaccct tccacgcatc agcat	25
<210> 114 <211> 25 <212> DNA	
<213> Artificial Sequence	
<220> <223> Description of Artificial Sequence: Primer	
<400> 114 cctccaggaa gcctgcgccg agaag	25
<210> 115 <211> 26 <212> DNA	
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<220> <223> Description of Artificial Sequence: Primer	
<400> 115 ggacattgtc cgtgatcagc aatcga	26

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<210> 116
<211> 25
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 116
                                                                   25
cagcetetgg aacaaccaga cgctg
<210> 117
<211> 25
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 117
                                                                   25
gtcaccgcga ggtacaagcc accac
<210> 118
<211> 25
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 118
                                                                   25
gcagctctgg agctctgtac cacct
<210> 119
<211> 25
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 119
acggccacgt cgagaatctg agcaa
                                                                   25
<210> 120
<211> 25
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 120
cgaagtgctc gcaagcaatg ccgaa
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<210> 121
<211> 35
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Primer
<400> 121
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atcccgggcg gtcgagtcgt attaggtgtt gtttc
<210> 122
<211> 30
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 122
                                                                   30
gageteeggt aggteegace tetteaattg
<210> 123
<211> 26
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 123
                                                                   26
gggcaactgt caatagcaga cctgga
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<211> 26
<212> DNA
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<223> Description of Artificial Sequence: Primer
<400> 124
                                                                   26
gcaagtccca acgaacgtgt ctcgct
<210> 125
<211> 25
<212> DNA
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<400> 125 gcgaagatga cgactgctat tgcga	25
<210> 126 <211> 25 <212> DNA <213> Artificial Sequence	
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<400> 126 cgtgatgact ccaatgctcc atacg	25
<210> 127 <211> 27 <212> DNA <213> Artificial Sequence	
<220> <223> Description of Artificial Sequence: Primer	
<400> 127 gccagcatcg aggtcagtat ccggtgt	27
<210> 128 <211> 27 <212> DNA <213> Artificial Sequence	
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<400> 128 gtctgtggcc ttcagaggcg catcctc	27